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/LEADSTALLON-TYPERRYRREHRPRSHLGQILRRRPWLVHPRHRYRWRRKNGI
FNTRLSRTFGYTVKRTTVTTPSWAVDWMRFNINDFVPPGGGTNKISIPFEYYRIRKVK
VERPWCCSPTITQCDRGVGSTAVTLDDNFVTRATALTYDPYVNYSSRHTTPQPFSYHSRY
FTPRPVLDSTIDYGONNKRNQLWLRLQTSRNVDHVGLGTAFENSKYDQDXNIRVTMY
VQFREFNLKDPPLNP
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Pred. No. 5.6e-168;
0; Mismatches 49;
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                            /protein_id="AAL69969;
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VGEGONEGGRPPHLGGFPNPVKKGPFNKVKWYTGARCHIREKAKGTDQONKBYCSKEGN
LLIECGAPRSQGORSDLSTAVSTLLESGSLYTVAEQHPVTFVRNFRGLELLKVSGKM
ÇKRDMKTNVHVIVGPPGCGKSKWAARFADFPETTYWRPPRNKWMDGYHGEBVVJDDFY
GWLPWDDLLRLCORYPLTVETKGGTVPFLPSSMLITSNQTPLEWYSSTAVPAVEALYR
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                                                                      GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCC
                                                      AGTGCTGTTATTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC
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Direct Submission
Submitted (02-JAN-2002) Hog Cholera Department, National Instifor Animal Health, 376 Chung-cheng Road, Tamsui, Taipei 25101,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Complete nucleotide sequences of porcine circovirus type
in plgs with various clinical syndromes
Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                   circovirus type 2 strain SC, complete genome
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porcine circovirus type 2.
Viruses; ssDNA viruses; Circoviridae; Circovirus.
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/protein_id-"AAL69968.1"
/db_xref-"GI:18448943"
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/db_xref="taxon:85708"
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51. .995
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Porcine circovirus type
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porcine circovirus type 2.

porcine circovirus type 2.

Viruses; ssDNA viruses; Circoviridae; Circovirus.

1 (bases 1 to 1767)

Mankertz,A., Domingo,M., Folch,J.M., LeCann,P., Jestin,A.,

Segales,J., Chmielewicz,B., Plana-Duran,J. and Soike,D.

Characterisation of PV-2 isolates from Spain, Germany and France
Virus Res. 66 (1), 65-77 (2000)
                                                                                                                                         2 (bases 1 to 1767)
Mankertz,A., Domingo,M., Folch,J.M., LeCann,P., Jestin,A., Segalas,J., Chmielewicz,B., Plana-Duran,J. and Soike,D.
Direct Submission
Submitted (03-NOV-1999) P24, Robert Koch Institut, Nordufer Berlin 13353, Germany
                                                                                                                                                                                                                                           /organism="porcine circovirus /isolate="FRA3"
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/gene="cap"
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                                    Circovirus sequences related to piglet weight loss disease (pwd) Patent: Wo 9929871.A 10 17-JUN-1999; HUTET EVELYNE (FR): ALBINA EMMANUEL (FR)
Location/Qualifiers
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Viruses; ssDNA viruses; Circoviridae; Circovirus 1 (bases 1 to 1767)
Hutet, E. and Albina, E.
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larity 100.0%; Pred. No. 1.7e-190;
Conservative 0; Mismatches 0;
                                                                                                   /organism="porcine circovirus"
/db_xref="taxon:46221"
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Matches 702; Conser
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/do_xref="G1:7021362"
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VEFWPCSPITQGDRGVGSSAVILDDNFVTKATALTYDPYVNYSSRHTITQPFSYHSRY
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VQFREFNFKDPPLNP"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                               /product="putative capsid protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
.737)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="replicase"
complement(36.
                                                                                                                                        /codon_start=1
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821. .1765
/gene="rep"
/note="Rep"
                                                /gene="cap"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 2000 nes 700; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           617
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circular VRL 23-FEB-2000

AF201311 1767 bp DNA circular VRL Porcine circovirus type 2 isolate FRA3, complete genome. AF201311 GI:7021361

AF201311/C LOCUS DEFINITION ACCESSION VERSION

gene 821. 1765 CDS /gene="rep" 821. 1765 CDS /gene="rep" /codon_start=1 /product="replicase" /product="replicase	Query Match 99.5%; Score 698.8; DB 14; Length 1767; Best Local Similarity 99.7%; Pred; No. 1.4e-189; Matches 700; Conservative 0; Mismatches 2; Indels 0; Gaps 0; QY 1 ATGACGTATCCAAGGGGGTTACCGAAGAAGAAGACACGCCCCCCCC	181 ACAGTCAGAACGCCTCCTGGGCGGTGGACATGATGATTTAATTAA	Db 377 AGTGCTGTTATTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC 318	QY 601 GCGTTCGAAAACAGTATATACGACCAGGAATACCATATACCATGTATGT
241 CCCCAGGAGGGGTCAAACCCCGGTGGACATGATGATTAATGACTTTCTT 498 241 CCCCAGGAGGGGGTCAAACCCCCGCTCTGGCCCTTTGAATACTACAGATAAGAAG 300 1111111111111111111111111111111111	317 TATGTAAACTACTCCCGCCATACCAGACCCTTCTCCTACCACCCGGTAC 258 481 TTTACCCCCAAACTGTCCTAGATTTCACTATGATTACTTCCAACCAA	36 Inear VRL Omplete geno Irus.	Isolation and Identification of Upublished 2 (bases I to 1767) 2 (Li,S.J., Li,Y., Li,J.W., Jin,H Direct Submission Submitted (14-UUN-2002) Harbin CAAS, No. 427, Maduanjie, Nangar 150001, P. R. China 1. 1767 2. 1. 1767 3. 1. 1767 3. 1. 1767 4. 176	<pre>gene</pre>

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                                                                                                                                                                                                                                                                                                                                                                                                          complement(989. .1033)
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                                                                                                                                                                            complement(688. .753)
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                                                                                                                                      complement(688. .753)
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/gene="ORF6"
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/gene="ORF10"
/note="4.1 kDa"
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RITSLVFWKNATEQSTEEGGGFVTLSPPCPEFPYEINY"
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                                                                                     porcine circovirus type 2.

porcine circovirus type 2.

viruses; sabNA viruses; Circoviridae; Circovirus.

1 (bases 1 to 170.)

Wellenberg, G.J., Pesch, S., Berndsen, F.W., Steverink, P.J.G.M., Hunneman, W., Van der Vorst, T.J.K., Peperkamp, N.H.M.T., Ohlinger, V.F., Schippers, R., Van Oirschot, J.T. and de Jong, M.F. Isolation and characterization of porcine circovirus type 2 from pigs showing signs of post-weaning multisystemic wasting syndrome in The Netherlands
                                                                                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (04-NOV-1999) Virology and Molecular Epidemiology,
BloScreen European Veterinary Disease Management Center GmbH, 11
Mendelstr., Muenster, NRW 48149, Germany
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note-"putative Rep protein; 35.8 kDa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          complement(join(1731. .1767,1. .92))
/gene="ORE9"
/note="4.6 kDa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complement(join(1731. .1767,1. .92))
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Porcine circovirus type 2, complete genome.
AF201897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /evidence=not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .565)
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                                                 AF201897.1 GI:11907587
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                                                                                                                                                                                                                                                                                                    source
        DEFINITION
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VERSION
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3 (bases 1 Meehan, B.M.
                                                                                                  Unpublished
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Matches 690; Conserv
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                                                                       Gaps
                                                                                                                                                                                                                                                                             Viruses; ssDNA viruses; Circoviridae; Circovirus.
1 (bases i to 1766)
Meehan, B.M. MCMailly, F., McNair, I., Walker, I., Ellis, J.A.,
Krakowka, S. and Allan, G.M.
                                                     1767;
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                 /translation="MAAGAVSSSPVTPPWIRHI" 361 c 500 g 458 t
                                                     690.8; DB 14;
No. 2.7e-187;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein
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                                                                       0; Mismatches
/protein_id="AAG41232.1"
/db_xref="GI:11907594"
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                                                     Score Pred. 1
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/translation="MPSKKNGRSGPQPHKRWVFTLNNPSEDERKKIRDLPISLFDYFI
VGEGRREGREPPHLOGFARNFVKKQTFRKVKWMTGARGHIERARGTDQOWREYCSKEGN
LLIEGGAPRSGGQRSFPPHLOGFARNSKKQTFVARDGRPUTFVRNFRGLAELLKVSGKM
QKRDWYTNVHVIVGPPGGGKSKWAANFADPETTYWKPPRNKWMDGYHGEEVVVIDDFY
GWLPWDDLLRLCDRYPLTVEFKGGTVPFLARSILITSWQTPLEWXSSTAVPAVEALYR
STISLYPWKNATEQSTEGGY
363 G 457 L
363 G 457 L
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KINGDOM
Isolation and characterization of porcine circovirus 2 from cask of sow abortion and porcine dermatitis and nephropathy syndrome Arch. Virol. 146 (4), 835-842 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 ATGACGTATCCAAGGAGGCGTTACCGAAGAAGAAGACACCGCCCCCCCGCAGCCATCTTGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      porcine dermatitis
                                                                                                                              Meehan, B.M., McNeilly, F., NcNair, I., Walker, I., West, K., Neehan, B.M., Krakowka, S., Kennedy, S. and Allan, G.M. Isolation and characterization of porcine circovirus type 2 fraces of sow abortion and porcine dermatitis and nephropathy
                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (25-AUG-2000) Meehan B.M., Veterinary Science,
Queen's University of Belfast, Stormont, BT4 3SD, UNITED
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1766;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein'
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Pred. No. 5.5e-185;
); Mismatches 12;
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/product="replication-associated
/protein_id="CAC41085.1"
/db_xref="G1:14330691"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:4621"
/db_xref="taxon:46221"
/country="United Ringdom:UK"
/note="pCV2 isolated from case onephropathy syndrome in the UK"
59. 1003
                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="porcine circovirus"
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59. .1003
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/product="unknown"
/protein_id="AAK60465.1"
/db_xref="G1:1442169"
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/db_xref="G1:14421167"
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TLLHFPAHFOKFSOPAEISDKRYRVLLCNGHQTTALQOGTHSTRQVTPLSLRSRSSTL
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/db_xref="G1:14422170"
/translation="WFIIHIGLSGGSLRLNSLNCTYMVTRIL"
complement(1022. 1027)
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/translation="MASSTPASPAPSDILSRKPQSERPPGRWT"
1523. 1630
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complement(1521. 1610)
/note="ORF6"
/note="glycosylation site"
906. .914
                                           /note="glycosylation site"
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/note="glycosylation site"
complement(688. .753)
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/note="ORF3"
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                                                                                                                                                     /product="unknown"
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/note="ORF10"
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/note="ORF5"
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/note="ORF7"
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                                                                   TATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTAC 1263
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VGEEGNEEGRTPHLQGFANFVKKQTFNKVKWYFGARCHIEKAKGTDQQNKEYCSKEGN
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QKRDWKTNVHVIVGPPGCGKSKWAANFADPETTYWKPPRNKWWDGYHGEEVVVIDDFY
GWLPWDDLLRLCDRYPLTVETKGGTVPFLARSILITSNQTPLEWYSSTAVPAVEALYR
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join(1749. 1767,1. 13)
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porcine circovirus type 2.

Viruses; ssDNA viruses; Circoviridae; Circovirus.

1 (bases 1 to 1767)

Shengbo,C. and Huanchun,C.

Cloning and sequence analysis of the genome of Porcine Circovirus type 2 isolated from pig with PMMS in China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (16-MAY-2001) Animal Medicine, Hua Zhong Agricultural University, Shi 21 Shan, Wuhan, Hubei 430070, P. R. China Location/Qualifiers
                                           TATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTAC
                                                                                                                             541 AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
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/country="China"
/note="isolated from a pig with PMWS"
complement(join(1731. 1767,1. 92))
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Porcine circovirus type 2, complete genome.
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816. .824
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/db_xref="GI:14422174"
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/protein_id="AAK60462.1"
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Shengbo, C. and Huanchun, C.
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porcine circovirus type 2
Viruses; ssDNA viruses; Circoviridae; Circovirus.
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                                                                     . 643.4; DB 14;
NO. 1.2e-173;
smatches 36;
                                  /translation="MAAGAVSSSAETPPWIRHR"
363 c 498 g 451 t .
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SPA2,
/evidence=not_experimental
/product="unknown"
/protein_id="AAK60468.1"
/db_xref="GI:14422172"
                                                                                     0; Mismatches
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/translation="MPSKKNGRSGPQPHKRWVFTLNNPSEDERKKIRELPISLFDYFI
VGEGERREGGRTPHLQGFANFVKKUFTRVKWYLGARCHIERAGTODGWREVCSKBGN
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OKRDWKTNVHVIVGPPGGGKSKWAANFADPETTYWKPPRNKWWDGYHGEEVVVIDDFY
GWLPWDDLLRLCDRYPLTVETKGGTVPFLARSILITSWQTPLEWYSSTAVPAVEALYR
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                                                                                                                                                                           Robert Koch Institut, Nordufer
                                                                                           2 (bases 1 to 1768)
Mankertz,A., Domingo,M., Folch,J.M., LeCann,P., Jestin,A.
Segales,J., Chmielewicz,B., Plana-Duran,J. and Soike,D.
Direct Submission
Submitted (03-NOV-1999) P24, Robert Koch Institut, Norduf
Berlin 13353, Germany
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 Soike, D.
Germany
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Chmielewicz, B., Plana-Duran, J. and ution of PCV-2 isolates from Spain, 66 (1), 65-77 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                           /product="putative capsid protein"
/protein_id="AAF35301.1"
/db_xref="G1:7021357"
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Pred. No. 6.8e-169;
0; Mismatches 47;
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/isolate="spac"
//db_xref="taxon:85708"
//db_xref="taxon:85708"
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/db_xref="G1:7021356"
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/gene="cap"
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822. 1766
/gene="rep"
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Best Local Similarity 93.3%;
Matches 655; Conservative
Segales, J., Chmielew
Characterisation of
Virus Res. 66 (1), 6
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porcine circovirus type 2-C
viruses; ssDNA viruses; Circoviridae; Circovirus.
1 (bases 1 to 176)
Hamel, A.L., Lin, L.L., Sachvie, C., Grudeski, E. and Nayar, G.P.
PCR detection and characterization of type-2 porcine circovirus
Can. J. Vet. Res. 64 (1), 44-52 (2000)
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                                                       Score 623.6; I
Pred. No. 5.6e·
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VGEGGNEEGRTPHLQGFANFVKKQTFNKVKWYLGARCHIEKAKGTDQONKEYCSKEGN
LLIECGAPRSOGORSDLSTANSTLLESGILVTVAKOHPVTFVKNFRGLAELLKVSGKM
GNRDWRTNVHFIVGPPGGKSKWAAANFANPETTYKKPPKNKWWDGYHGEKVVVIDDFY
GWRDWADDLLLALCDRYPELTVKTKGGTVPFLARSILITSNQTPLEWYSSTAVPAVEALYR
RITSLVFWKNATEQSTEEGGQFVTLSPPCPFLARSILITSNQTPLEWYSSTAVPAVEALYR
COMPLEMENL(1034. .1735)
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FFFRLSRTFGYTVKFTYTYTPSWAVDWHRFKIDDFVPPGGGTRKISIPFEYYRIRVY
VEFWPCSPTTGGDRGVATYVILDHPVTKATALTYDPYVNYSRHTIPQPFSYHSRY
FTPRPVLDSTIDYFQPNNKRNDLWARLGTSGNVDHVGLGTAFBNSKYDQDYNIRVTWY
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AACCAGCTGTGGCTGACACTACAAACTGCTGGAAATGTAGAACCACGTAGGCCTCGGCACT
                                                                                                                                                                                                                                                                                      AGTGCTGTTATTTTAGATGATAACTTTGTAACAAAGGCCACACACCCTCACCTATGACCCC
                                                                                                                                                                AATCAGCTTTGGCTGAGGCTACAAACCTCTGCAAATGTGGACCACGTAGGCCTCGGCACT
                                                                                                                                                                                                                                                                          of Saskatchewan,
S7N 5E3, Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wang, L., Willson, P., Chow, B., Gibbons, E. and Babiuk, L. Direct Submission Submitted (26-AUG-1998) VIDO, University of Saskatchew Veterinary Road, Saskatcon, Saskatchewan S7N 5E3, Canal Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gibbons, E. and Babiuk, L.
                                                                                                                                                                                                                                                                                                                               TTCAGAGAATTTAAATTTTAAAGACCCCCCCCTTAACCCTTAA 702
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                                                                                                                                                                                                                                                                                                                                           /organism="porcine circovirus"/strain="9741"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Viruses; ssDNA viruses; Circoviridae;
1 (bases 1 to 1768)
Wang, L., Willson, P., Chow, B., Gibbons,
Emergence of a new porcine circovirus
Unpublished
2 (bases 1 to 1768)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /protein_id="AAC61739.1"
/db_xref="G1:3661519"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /protein_id="AAC61740.1"
/db_xref="GI:3661520"
                                                                                                                                                                                                                                                                                                                                                                                                                     Porcine circovirus strain 9741, AF086835.1 GI:3661518
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/translation="MTYPRRRYRRRYRPSHLGQILRRRPWLVHPRHRYRWRKNGI
TYMLLGRFGCYTVKATTVRYFSWAVDMLRFKLDDFVPPGGGTNKISIPPEYYRIRKVK
VEFRPCSPTTQGDRGVGSSAVILDDNFVPKATALTYDPYVNYSSRHTITQPFSYHSR"
COMplement (1519. . 1611)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1735 ATGACGTATCCAAGGAGGCGTTACCGCAGAAGAAGACCGCCCCCGCAGCCATCTTGGC 1676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1675 CAGATCCTCCGCCCCCCGCCCCTGGCTCCACCCCGCCACCCGTTACCGCTGGAGAGG 1616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1495 CCCCCAGGAGGGGGACCAACAAAATCTCTATACCCTTTGAATACTACAGAATAAGAAGA 1436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /translation="MVFIIHLGFKWGVFKIKFSELYIHGYTDIVVLVVYTVFERSAEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACAGTCAGAACGCCCTCCTGGGCGGTGGACATGAATGAGATTCAATATTAATGACTTTCTT 240
                                                                                                                                                                             complement(1256, 1735)
/note="similar to Porcine circovirus ORF-2 encoded by the sequence presented in GenBank Accession Number AF027217"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAGATCCTCCGCCGCCGCCCTGGCTCGTCCACCCCGCCACCGTTACCGCTGGAGAAGG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121 AAAAATGGCATCTTCAACACCCGCCTCTCCCGCACCTTCGGATATACTGTCAAGCGAACC 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 CCCCCAGGAGGGGGTCAAACCCCCGCTCTGTGCCCTTTGAATACTACAGAATAAGAAAG 300
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1524. .1631
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/db_xref="G1:4106892"
/translation="MAAGAGPSSAVTPPWIRHS"
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                                                                                                                                                                                                                                                                                             /protein_id="AAD03062.1"
/db_xref="GI:4106887"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /protein_id="AAD03066.1"
/db_xref="G1:4106891"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /protein_id="AAD03068.1"
/db_xref="GI:4106893"
                                                                       /protein_id="AAD03065.1"
                                                                                                                                                           complement(1022. .1027)
                                                                                        /db_xref="GI:4106890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /product="ORF-10"
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/product="ORF-6"
                           /codon_start=1
/product="ORF-5"
                                                                                                                                                                                                                                                                        /product="ORF-2"
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VGEBGENEGRETPHLQGFANFVKKQFTRVKWW.TGARCHTERKAGTDQORKEYGSREGN
LLIQCGAPRSGQQREFTPLLEGSSLYTYAROHPYTFYRNFRGLAELLKVSGKM
OKRDWKTNVHVIVGPPGCGKSKWAANFADPETTYWKPPRNKWWGSYHGEEVYVIDDFY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /translation="MVTIPPLVSRWFPVCGFRVCKISSPFAFATPRWPHNDVYIRLPI
fllhfPAHFQKFSQPAEISDKRYRVLLCNGHQTPALQOGTHSSRQVTALSLRSRSSTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /protein_id="AAD03070.1"
/db_xref="GI:4106895"
/translation="MYTSLWGHLGVAKANGLLILQTRKPHTGNHLETSGGMVTMVKKW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          circovirus presented in GenBank Accession Numbers AF027217 and U49186"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GWLPWDDLLRLCDRYPLTVETKGGPVPFLARSILITSNQTPLEWYSSTAVPAVEALYR
RITSLVFWKNATEQSTEEGGGFVTLSPPCPEFPYEINY"
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                                                                                                          3 (bases I to 1768)
Hamel, A.L. and Nayar, G.P.S.
Direct Submission
Submitted (27-NOV-1998) Virology Laboratory, Veterinary Diagnostic
Laboratory, Manitoba Agriculture, 545 University Crescent,
Winnipeg, Manitoba R3T 5S6, Canada
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                   using DNA extracted from lung, mesenteric lymph node and ornsil of pig, similar to Porcine circovirus sequence presented in GenBank Accession Number AF027217; type-C designation is based upon restriction endonuclease
                                                                                                                                                                                                                                                                                                                                                              /note="sequence obtained from several overlapping PCRs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            join(1762. 1768,1. .2)
/note="similar to the nonanucleotide motif of Porcine
                       nucleotide sequence of four different isolates of circovirus detected in pigs with various clinical syndromes unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="putative Rep and coat protein"
/protein id="AAD03061.1"
                                                                                                                                                                                                                                                                                             /organism="porcine circovirus type 2-C'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        digestion pattern"
join(1750, 1768,1, 13)
/note="putative replication site"
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/db_xref="GI:4106888"
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/db_xref="GI:4106889"
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complement(989. .1033)
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2 (bases 1 to 1768)
Hamel, A.L. and Nayar, G.P.S.
Nucleotide sequence of four
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/rpt_unit=13..18
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Institute of Technology, 300 Shei Fu Road, Lu Liao Li, Chia-Yi City
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Yang, K.H., Lee, Y.F., Chao, D.S., Shieh, Y.C. and Lai, S.S.
Complete nucleotide sequences of porcine circovirus Tainan
outbreak in Taiwan
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Yang, K.H., Lee, Y.F., Chao, D.S., Shieh, Y.C. and Lai, S.S.
Direct Submission
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porcine circovirus
Viruses; ssDNA viruses; Circoviridae; Circovirus.
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/protein_id="AAD50432.1"
/db_xref="GI:5739339"
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/organism="porcine circovirus"
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Porcine circovirus complete genome.
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/country="Taiwan"
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VGEGONEEGRPHLOGFANFVKKOTFNKVKWYFGARCHIEKAKGTDQQNKEYCSKEGN
LLIECGAPRASOGORSDLSTAVSTLLESSGSLVTVYAEQHPYFKNFRGLAELLKVSGKM
QKRDWKTNVHYLVGPPGGGKSKWAANFADPETTYWKPPRNKWNDGYHGEEVVYIDDFY
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                                                                1135 ATGACGTATCCAAGGAGGCGTTTCCGCAGACGAAGACACCGCCCCCGCAGCCATCTTGGC
                                            AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
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Viruses; ssDNA viruses; Circoviridae; Circovirus.
1 (bases 1 to 1768)
Kuo,T.Y., Chiou,Y.C. and Lai,S.S.
Complete nucleotide sequences analysis of porcine circovirus outbreak in Taiwan
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/isolare="MIW98"
/db_xref="taxon:46221"
/country="Taiwan"
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/db_xref="G1:5052005"
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/translation="MPSKKNGRSGPQPHKRWVFTLNNPSEDERKKIRELPISLFDYFI
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LLIECGAPRSGGQRSTPPHLGSGSLVYVABOHPVTFVRNFGLAELLKVSGKB
ULIECGAPRSGGGRSTPHLESGSLVTVABOHPVTFVRNFGLAELLKVSGKB
QKRDWKTNVHVIVGPPGCGKSKWAANFADPETTYWKPPRNKWWDGYHGEEVVVIDDFY
GWLEWNDLLRLCDRYPLTYSTKGGTVPFLARSTLITSKQTPLEWYSSTRVPAVBALRR
TTSLVFWRNATEGGFTEGGFFYLLSPRGTTYN"
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                                                                                                              Score 623.6; DB 14
Pred. No. 5.6e-168;
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circular VRL 23-FEB-2000

1768 bp DNA circular VRL 2 isolate GER3, complete genome.

Porcine circovirus type AF201307

AF201307

GI:7021349

AF201307.1

ACCESSION VERSION

porcine circovirus type 2.
porcine circovirus type 2
porcine circovirus type 2
Viruses; ssDNA viruses; Circovirus
1 (bases 1 to 1768)

ORGANISM

KEYWORDS SOURCE

REFERENCE

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/translation="MPSKKNGRSGPQPHKRWVFTLNNPSEDERKKIRELPISLFDYFI
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QKRDWKTNVHYIVOPPGCGKSKWAANFADPETTYWKPPRNKWMDGYHGEBTVVIDDFY
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FTPKPVLDRTIDYFQPROVGSTAYLILDNEVTKATALTYDYPVNYSSRHTIPOPFGYHSRY
FTPKPVLDRTIDYFQPROVISKNOLMARLQTSANVDHVGLGTAFFNSKYDODYNIRVTMY
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  , Jestin, A.,
Soike, D.
Germany and B
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Mankertz, A., Domingo, M., Folch, J.M., LeCann, P., Segales, J., Chmielewicz, B., Plana-Duran, J. and Scharacterisation of PCV-2 isolates from Spain, C Virus Res. 66 (1), 65-77 (2000)
                                                                                                                 2 (bases 1 to 1768)
Mankertz,A., Domingo,M., Folch,J.M., LeCann,P.,
Segales,J., Chmielewicz,B., Plana-Duran,J. and Edinect Submission
Submitted (03-NOV-1999) P24, Robert Koch Instituted in 13353, Germany
Location/Qualifiers
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/product="putative cap/
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/db_xref="taxon:85708'
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/gene="cap"
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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd.

May 18, 2003, 09:15:38 ; Search time 2022 Seconds (without alignments) 10103.941 Million cell updates/sec 702 1 atgacgtatccaaggaggcg.....accccccacttaacccttaa 702 4109280 Total number of hits satisfying chosen parameters: 2054640 seqs, 14551402878 residues OM nucleic - nucleic search, using sw model Post-processing: Minimum Match 08
Maximum Match 1008
Listing first 45 summaries IDENTITY_NUC Gapop 10.0 , Gapext 1.0 em_htg_other: Minimum DB seq length: 0 Maximum DB seq length: 200000000 US-09-514-245B-25 em_htg_hum:* em_htg_inv:* em_ro:* em_sts: gb_ro:* gb_sts: gb_htg:* gb_in:* gb_om:* gb_ov:* gb_pat: *
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Circovirus sequences related to piglet weight loss
Patent: WO 9929871-A 9 17-JUN-1999;
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Viruses; ssDNA viruses; Circoviridae; Circovirus.
1 (bases 1 to 1767)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTCAGAGAATTTAAATTTTAAAGACCCCCCACTTAACCCTTAA
                                          B-ORF2"
                                                                             100.0%; Score 702; DB 6; I
llarity 100.0%; Pred. No. 1.5e-190;
Conservative 0; Mismatches 0.
                     /organism="porcine circovirus"
/db_xref="taxon:46221"
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                                          type 1
153 t
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WO9929871.
                                         /note="Circovirus MAP
218 c 138 g
Location/Qualifiers
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Sequence 9 from Patent NAX003274
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        porcine circovirus.
                                                                                           Local Similarity
nes 702; Conserv
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Gaps

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1555

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1495

300

360

24 - AUG - 2000

PAT

RESULT 2 AX003274/C LOCUS DEFINITION

g 5

ORGANISM

REFERENCE AUTHORS

ACCESSION VERSION KEYWORDS SOURCE

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents the nucleotide sequence of PCV isolate Impl011-48285. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PWMS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from it, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific antibodies immunication, formal in any many in any i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG
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                                                                                                                                                                                                                                                                                                                                           Ellis J;
porcine multisystemic wasting syndrome; pig; vaccine; ss.
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                                                                                                                                                                                                                                                                                                                                           Charreyre CE, Clark E,
Hassard L, Meehan B;
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                                                                                                                                                                                             98FR-0000873
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                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-246948/21
                                                 Porcine circovirus.
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Haines D,
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Search completed: May 18, 2003, 11:07:55 Job time : 262 secs

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ВP
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                                     Ouery Match
Best Local Similarity 99.0%;
Matches 384; Conservative
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(UYBE-) UNIV QUEENS BELFAST
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                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine; pig;
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Harding J,
Mcneilly F;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention provides a novel antigenic preparation directed against porcine multisystemic wasting syndrome (PMWS) that comprises porcine circovirus (PCV) antigen and porcine parvovirus (PPV) antigen. The PCV antigens and PPV antigens can be used for vaccination against PPWS. The present sequence represents the DNA sequence of PCV Imp.1011-48121
                                                                                                                                              CTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTTT
                                                              AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC
                                                                                                 CTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGGTACTTTACCCCCAAACC
                                                                                                                                    TGTCCTAGATTTCACTATTGATTACTTCCAACCAACAACAAAAGAAACCAGCTGTGGCT
                                                                                                                                                                       GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of a porcine circovirus antigen and a porcine parvovirus antigen vaccination against porcine multisystemic wasting syndrome \,
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           Indels
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Pred. No. 5.6e-105;
0; Mismatches 4;
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99.08;
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           Conservative
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Best Local Similarity
Matches 384; Conserv
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1767 CTGGCCCTGCTCCCCGGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTCT 1708
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                                                                                             Gaps
                                                                                                                                                                Hassard L;
Audonnet JF;
                                            Length 1767;
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                                                                                             Indels
Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;
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GS,
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Krakowka (
                                         Score 381.6; DB 21;
Pred. No. 5.6e-105;
                                                                                           0; Mismatches
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Charreyre CE, Chappuis GE, K
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AAX35211
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                                                                                                                                                                                                                                              494
                                            The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment of a strain of PCV, which was used in the present invention.
                                                                                                                                                                                                                                                                                                                                                           315 CTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTTT
                                                                                                                                                                                                          AGATGATAACTTTGTAACAAAGGCCACACACCCTCACCTATGACCCTTATGTAAACTACTC
                                                                                                                                                                                                                                              CTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTACTTTACCCCCAAACC
                                                                                                                                                                                                                                                                                  GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG
                                                                                                                                DB 22; Length 1767;
                                                                                                                                                   ;
0
post-weaning, multisystemic wasting syndrome PCV-2
                                                                                                                                                   Indels
                                                                                                             Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  porcine multisystemic wasting syndrome; pig; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        porcine circovirus; PCV; PMWS;
                                                                                                                                1 Similarity 99.0%; Pred. No. 5.6e-105; 94; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of PCV isolate Imp1011-48285
                                                                                                                                                                                                                                                                                                                                                                                                           1407 TCTTAAAGACCCCCACTTAACCCTTAA 1380
                                                                                                                                                                                                                                                                                                                                                                                              675 ITTTAAAGACCCCCCACTTAACCCTTAA 702
                            Disclosure; Fig. 1 #2; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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AAX35379 standard; DNA; 1767
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98FR-0000873
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Porcine circovirus
and/or
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22-JAN-1998;
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Matches 384;
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The present sequence represents the nucleotide sequence of PCV isolate Impl011-48285. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PWMS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from lt, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific antibodies in any constant and any constant of the polypeptides can be used to detect antigens, in any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               315 CTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           555 GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   435 CTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTACTTTACCCCCAAACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ); DB 20;
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                                                                                              Clark E,
                                                                                              Charreyre CE, Clark
Hassard L, Meehan B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 378.4; I
Pred. No. 5.2e-
0; Mismatches
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                                                                                                                                                                                                                                                  New type II porcine circovirus
                                                                                                                                                                                                                                                                                                              Claim 11; Fig 2; 56pp; French.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 98.5%;
Matches 382; Conservative
(UYBE-) UNIV QUEENS BELFAST. (UYSA-) UNIV SASKATCHEWAN.
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                                                                                           Chappuis GE,
Harding J,
                                                                                                                                                                                      WPI; 1999-264024/22
                                                                                        Allan G,
Haines D,
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ATGTACAATTCAGAGAATTTAAATTTTAAAGACCCCCCACTTAACCTTAA 702
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ب
                                                                                                             PCV isolate; type II porcine circovirus; PCV; PMWS;
porcine multisystemic wasting syndrome; pig; vaccine; ss.
                                                                                                                                                                                                                                                         Ellis
                                                                                              Nucleotide sequence of PCV isolate Imp1011-48121.
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Hassard L, Me
                                                   BP
                                                                                                                                                                                                                                                                                            New type II porcine circovirus
                                                                                                                                                                                                                                                                                                          Claim 11; Fig 1; 56pp; French.
                                            378/c
AAX35378 standard; DNA; 1767
                                                                                                                                                                                                    97FR-0012382.
98FR-0000873.
                                                                                                                                                                                                                                 (UYBE-) UNIV QUEENS BELFAST. (UYSA-) UNIV SASKATCHEWAN.
                                                                                                                                                                                              98FR-0003707
                                                                                                                                                                               98WO-FR02107
                                                                               (first entry)
                                                                                                                                                                                                                                                        Chappuis GE,
Harding J,
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                                                                                                                                   Porcine circovirus
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(UYBE-) UNIV QU
                                                                                                            PCV isolate;
                                                                                                                                                W09918214-A1
                                                                                                                                                                               01-OCT-1998;
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22-JAN-1998;
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                                                                                                                                                                                                                                                                Haines D,
             664
                                                                 AAX35378;
653
                                                                                                                                                                                                                                                         Allan G,
                                    RESULT 10
                                            AAX35378/
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The present sequence represents the nucleotide sequence of PCV isolate Impl011-48121. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PWMS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from it, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific anthodies, while antibodies and a staised against the polypeptides can be used to detect antigens, in any usual immunoassay

Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;

Query Match

DB 20; Length 1767;

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1161 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 1111 | 11
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54.4%; Score 381.6; DB 20; 99.0%; Pred. No. 5.6e-105; ive 0; Mismatches 4;
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                                                                                                         Similarity
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Length 1767;

20;

DB

Score 381.6;

54.48;

Query Match

Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;

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The present sequence represents the nucleotide sequence of PCV isolate Impl011-48121. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PWWS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from it, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA polypeptides, also as probes or primers for diagnosts in usual hybridization or amplification assays. These polypeptides may also beused diagnostically to detect PCV-specific antibodies, while antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             raised against the polypeptides can be used to detect antigens, in any
                                                        555 GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAAAAGAG
                                                                                                                                                      TATATACGACCAGGAATACAATATCCGTGTAACCATGTACTACAATTCAGAGAATTTAA
                                         New type II porcine circovirus, used for, e.g. passive immunization of pregnant sows
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Hassard L, Meehan B;
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Harding J,
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Haines D,
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97US-0069233
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                                Potter
                   (UYSA-) UNIV
                                                         isolated
16-DEC-1997;
11-DEC-1997;
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                                                 GAGAAGGAAAAATGGCATCTTCAACACCCGCCTCTCCCGCACCTTCGGATATACTGTCAA
                                                                                                                     CTTTCTTCCCCCAGGAGGGGGGTCAAACCCCCGCTCTGTGCCCTTTGAATACTACAGAAT
                                                                                                                                              AAGAAAGGTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGT
                                                                                                                                                                       GGGCTCCAGTGCTGTTATTTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTA
                                                                                                                                                                                               TGACCCCTATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTC
                                                                                                                                                                                                                        CAAAAGAAACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCT
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                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                Porcine circovirus type II; PCVII; PCVI; pig; infection; vaccine; postweaning multisystemic wasting syndrome virus; diagnosis; ds.
                                                                                                                                                                                                                                                                                                        circovirus type II B9 nucleotide sequence fragment #2.
                                                                                                                                                                                                                                                                                                  ATGTACAATTCAGAGAATTTTAAATTTTAAAGACCCCCCACTTAACCCTTAA
ed. No. 8.6e-170;
Mismatches 49;
Pred.
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Local Similarity 92.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 590.8; DB 20;
Pred. No. 2.1e-168;
                                                                                                                                                                                                                                           present invention describes a new isolated
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Local Similarity 91.5%;
nes 650; Conservative
SASKATCHEWAN
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RESULT 7
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Charreyre CE, Chappuis GE, K
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AAF75841 standard; DNA; 1768
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(UYSA-) UNIV SASKATCHEWAN.
(UYBE-) UNIV QUEENS BELFAST.
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                                                                                                                                 The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat plgs against impocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequale associated with PCV-2. The present sequence is a DNA fragment of a strain of PCV, which was used in the present invention.
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Use of porcine circovirus-2 immunogen to formulate a vaccine composition to treat pigs against myocarditis, abortion, intrinfection and/or post-weaning, multisystemic wasting syndrome associated with PCV-2
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                                                                                                                                                                                                                                                                                        87.9%; Score 617.2; DB 22;
92.5%; Pred: No. 2.5e-176;
Mismatches 53;
                                                                                                    English
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nes 649; Conservative
                                                                                                Claim 11; Fig 7; 134pp;
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Type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAX83754 to AAX83757 represent PCVII nucleotide sequences. AAY24929 to AAY24934 represent PCVII open reading frame (ORF) proteins (N.B. the PCVII ORFs given in Fig 2A to Fig 2B do not correspond exactly with the PCVII ORFs given in Fig 3A to Fig 3D). The PCVII polypeptides can be used for treating or preventing PCVII infection in vertebrates. The products can also be used to detect the
                                                                                                   AGTGCTGTTATTTTAGATGATAACTTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC
                                                                                   TATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTAC
                                                                                                                                                      AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
                                                                                                                                                                                                                                         The present invention describes a new isolated porcine circovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Porcine circovirus type II; PCVII; PCVI; pig; infection; vaccine postweaning multisystemic wasting syndrome virus; diagnosis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 1786;
                                                                                                                                                                                                                                                                                                                                                                            1075 TTCAGGGAATTTAATCTTAAAGACCCCCCACTTAACCCCTAA 1034
                                                                                                                                                                                                                                                                                                                                                           661 TTCAGAGAATTTAAATTTTAAAGACCCCCCACTTAACCCTTAA 702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1786 BP; 467 A; 367 C; 481 G; 471 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Porcine circovirus type II 9741 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 595.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Willson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated porcine circovirus Type II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Fig 4; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX83755 standard; DNA; 1786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-CA01130.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0069750.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0069233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYSA-) UNIV SASKATCHEWAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Babiuk LA, Potter AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-394957/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Porcine circovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W09929717-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1556
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or sost-weaming, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment of a strain of PCV, which was used in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to the use of porcine circovirus-2 (PCV-2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Use of porcine circovirus-2 immunogen to formulate a vaccine composition to treat pigs against myocarditis, abortion, intrauterine infection and/or post-weaning, multisystemic wasting syndrome associated with PCV-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1615 AAAAATGGCATCTTCAACACCCGCCTCTCCCCCACCTTCGGATATACTGTCAAGCGTACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGCCTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAGATCCTCCGCCGCCGCCCTGGCTCGTCCACCCCGCCACCGTTACCGCTGGAGAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACAGTCAGAACGCCCTCCTGGGCGGTGGACATGATGATGATGATTATTAATGACTTTCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCCCCAGGAGGGGGGGTCAAACCCCCGGTCTGTGCCCTTTGAATACTACAGAATAAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Audonnet JF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22; Length 1768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
0
                                                                                                   Vaccine; pig; myocarditis; abortion; intrauterine infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 612.4; DB 22; Length
Pred. No. 7e-175;
0; Mismatches 56; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1768 BP; 450 A; 360 C; 496 G; 460 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                           , Haines D,
Krakowka GS,
                                                                                                                                                                                                                                                                                                                                                                                                             ы
                                                                                                                                                                                                                                                                                                                                                                                                          Allan GM, Meehan B, Clark E
Charreyre CE, 'Chappuis GE,
                                                                  PCV DNA fragment of Imp 1103 strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; Fig 6 #2; 134pp; English.
                                                                                                                   multisystemic wasting syndrome; ds
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92.0%;
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(UYSA-) UNIV SASKATCHEWAN.
(UYBE-) UNIV QUEENS BELFAST.
                                                                                                                                                                                                                                                       28-AUG-2000; 2000WO-EP08781
                                                                                                                                                                                                                                                                                       31-AUG-1999; 99US-0151564
31-MAY-2000; 2000US-0583350
                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  646; Conservative
                                                                                                                                                    Porcine circovirus-2.
                                                                                                                                                                                                                                                                                                                                                                                                             Allan GM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                     WO200116330-A2
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Mcneilly F;
                                 18-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                           Ellis JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121
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AAF75840;
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Best Local $
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1538 ACTGCTGTTATTCTAGATGATAACTTTGTAACAAGGCCACAGCCCTAACCTATGACCCA 1479
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                                                                                                                                                        CAGATCCTCCGCCGCCGCCCCTGGCTCGTCCACCCCGGCACCGTTACCGCTGGAGAGG 120
                                                                                                                                                                                                                                                                                                    CCCCCAGGAGGGGGGTCAAACCCCCGCTCTGTGCCCCTTTGAATACTACAGAATAAGAAAG
                                                                                                                                                                                                                                                                                                               ACAGTCAGAACGCCCTCCTGGGCGGTGGACATGAGATTCCAATATTAATGACTTTCTT
                                                                                                                                                                                                                                                                                                                                                                                                AGTGCTGTTATTTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                             TATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
                                     828 A; 455 C; 538 G; 948 T; 0 other;
                                                           Score 622; DB 22;
Pred. No. 1.1e-177;
                                                                                    0; Mismatches
              a plasmid, which expresses gene
  associated with PCV-2.
                                                           88.6%;
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                                                                                    652; Conservative
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                                     BP;
                                                                        Similarity
                                    Sequence 2769
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                                                             Query Match
                                                                          Local
                                                                                    Matches
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GTTAAGGTTGAATTCTGGCCCTGCTCCCCCATCACCCAGGGTGATAGGGGAGTGGGCTCC 1761
                                  2001
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                                                                                           1822 TATGTAAACTACTCCTCCGCCATACAATCCCCCAACCCTTCTCCTACCACTCCCGTTAC 1881
                                                                                                                     540
                                                                                                                                                                                                                            AGTGCTGTTATTTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or
                                                                                                                                                                             TATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTAC
                                                                                                                  AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
                                                                                                                                                                                                             Use of porcine circovirus-2 immunogen to formulate a vaccine composition to treat plus against myocarditis, abortion, intrauterine infection and/or post-weaning, multisystemic wasting syndrome associated with PCV-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JF;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ALVAC; porcine circovirus-2; vaccine; pig; myocarditis; abortion; intrauterine infection; multisystemic wasting syndrome; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hassard L;
Audonnet J
                                                                                                                                                                                                                                                                          TTCAGAGAATTTAAATTTTAAAGACCCCCCACTTAACCCTTAA 702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ellis JA, Allan GM, Meehan B, Clark E, Haines D,
Harding J, Charreyre CE, Chappuis GE, Krakowka GS,
Mcneilly F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3 #3; Fig 6 #1; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                    pJP107 DNA fragment
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(UYSA-) UNIV SASKATCHEWAN.
(UYBE-) UNIV QUEENS BELFAST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-2000; 2000WO-EP08781
                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-244408/25.
                                                                                                                                                                                                                                                                                                                                             AAF75830 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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                                             1762
                                                                                                                                          1882
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                      361
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                                                                                                                                                                                                                                                                                                                                                                     AAF75830;
                                                                                                                                                                                                                                                                                                                                                                                                                    Plasmid
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Gaps

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Indels

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Length 2769;

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PCV2; porcine circovirus 2; virucide; immunostimulant; vaccion postweaning multisystemic wasting syndrome; PMWS; infection; pig pathogen; open reading frame; ORF; ss.
              TTCAGAGAATTTAATCTTAAAGACCCCCCACTTAAACCCTAA 1197
TTCAGAGAATTTAAATTTTAAAGACCCCCCCACTTAACCCTTAA
                                                                                                                                                                                                                                pJP107 donor plasmid for PCV2 ORF2 and ORF1
                                                                                                                                                                                                                                                                                                                                           Porcine circovirus
                                                                                                  AAF28320/C
ID AAF28330/C
XX
AC AAF283
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DE DJP107
XW
PCV2;
KW
POStwe
XX
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post-weaning, multisystemic wasting syndrome and other pathological

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ATGACGTATCCAAGGAGGCGTTACCGCAGAAGAGACACCGCCCCCCCGCAGCCATCTTGGC 1676
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAX83754 to AAX83757 represent PCVII nucleotide sequences. AAX2429 to AAX2434 represent PCVII open reading frame (ORF) proteins (N.B. the PCVII ORFS given in Pig 2A to Fig 2B do not correspond exactly with the PCVII ORFS given in Fig 3A to Fig 3B). The PCVII polypeptides can be used for treating or preventing PCVII infection in vertebrates. The products can also be used to detect the
                           CAGATCCTCCGCCGCCCCCTGGCTCGTCCACCCCCCGCCATACCGCTGGAGAGG
  541 AACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a new isolated porcine circovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Porcine circovirus type II; PCVII; PCVI; pig; infection; vaccine; postweaning multisystemic wasting syndrome virus; diagnosis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                           1238 TTCAGAGAATTTAATCTTAAAGACCCCCCACTTAAAACCCTAA 1197
                                                                                                                                                                                               702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1768 BP; 454 A; 361 C; 492 G; 461 T; 0 other;
                                                                                                                                                                                                  TTCAGAGAATTTAAATTTTAAAGACCCCCCCACTTAACCCTTAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Porcine circovirus type II 412 nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 620.4; DB 20;
Pred. No. 2.6e-177;
0; Mismatches 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Willson
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                                                                                                                                                                                                                                                                                                                                                                        BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Fig 2; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wang
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92.78;
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Best Local Similarity 92.7
Matches 651; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Potter AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-394957/33
                                                                                                                                                                                                                                                                                                                                                                      AAX83754 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence was used in the construction of a recombinant virus comprising DNA from porcine circovirus 2 (PCV2). The recombinant virus is useful as vaccine for treatment and prophylaxis of PCV2 infection, such as postweaning multisystemic wasting syndrome (PWMS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9
                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel recombinant virus comprising DNA from porcine circovirus 2 usefu
as vaccine for treatment and prophylaxis of porcine circovirus
infection, such as postweaning multisystemic wasting syndrome in pigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1898 ATGACGTATCCAAGGAGCGTTACCGCAGAAGAAGACACCGCCCCCCGCAGCCATCTTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ATGACGTATCCAAGGAGGCGTTACCGAAGAAGAAGACACCGCCCCGCAGCCATCTTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAGATCCTCCGCCGCCCCTGGCTCGTCCACCCCCGCCACCGTTACCGCTGGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAATGGCATCTTCAACACCCGCCTCTCCCGCACCTTCGGATATACTGTCAAGCGAACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACAGTCAGAACGCCCTCCTGGGCGGTGGACATGATGAGATTCAATATTAATGACTTTCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCCCAGGAGGGGGTCAAACCCCCGCTCTGTGCCCTTTGAATACTACAGAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGTGCTGTTATTTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3609 BP; 1050 A; 618 C; 737 G; 1204 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 622; DB 22;
Pred. No. 1.2e-177;
); Mismatches 50;
                                                                                                                                                                                                                                                                                                                      Charreyre CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Fig 6; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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92.9%;
                                                                                                                                               2000WO-IB00882.
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                                                                                                                                                                                                                                                                                                                      Perez JM,
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-080692/09
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Local Similarity
nes 652; Conserv
                                               WO200077216-A2.
                                                                                                                                                                                                                                                                        (MERI-) MERIAL
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                                                                                                                                                                                                                      01-JUN-2000;
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                                                                                                                                                                                             10-JUN-1999;
                                                                                               21-DEC-2000
Synthetic.
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1998.
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/SIDS2/gagdata/geneseq/geneseqn-embl/NA1984.
/SIDS2/gagdata/geneseq/geneseqn-embl/NA1985.
/SIDS2/gagdata/geneseq/geneseqn-embl/NA1986.
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GenCore version 5.1.6
Copyright (c) 1993 • 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2185239 segs, 1125999159 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                        OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                   IDENTITY_NUC Gapox 10.0
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                              Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Searched:
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                                                                                                                                                                                     Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description.	Plasmid pJP102 DNA	pJP102 donor plasm	Plasmid pJP107 DNA	pJP107 donor plasm	Porcine circovirus	PCV DNA fragment o	PCV DNA fragment o	Porcine circovirus	Porcine circovirus
SUMMARIES	QI	AAF75827	AAF28317	AAF75830	AAF28320	AAX83754	AAF75841	AAF75840	AAX83755	AAX83757
	,	22	22	22	22	20	22	22	20	20
	Match Length DB	2520	2520	2769	3609	1768	1768	1768	1786	1361
ع مو	Match	88.6	88.6	88.6	88.6	88.4	87.9	87.2	84.8	84.2
	Score	622	622	622	622	620.4	617.2	612.4	595.6	590.8
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ALIGNMENTS

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Hassard L;
Audonnet JF;
                                                                                                             ALVAC; porcine circovirus-2; vaccine; pig; myocarditis; abortion; intrauterine infection; multisystemic wasting syndrome; ds.
                                                                                                                                                                                                                                                                                                                   Ellis JA, Allan GM, Meehan B, Clark E, Haines D, F
Harding J, Charreyre CE, Chappuis GE, Krakowka GS,
Mcneilly F;
                     AAF75827 standard; DNA; 2520 BP.
                                                                                                                                                                                                                                                                        (MERI-) MERIAL.
(UYSA-) UNIV SASKATCHEWAN.
(UYBE-) UNIV QUEENS BELFAST.
                                                                                                                                                                                                                                       31-AUG-1999; 99US-0151564.
                                                                                        Plasmid pJP102 DNA fragment
                                                                                                                                                                                                                28-AUG-2000; 2000WO-EP08781
                                                                 18-MAY-2001 (first entry)
                                                                                                                                                                     WO200116330-A2
                                                                                                                                               Unidentified.
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                                           AAF75827;
RESULT 1
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                                                                                                                           immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment of a plasmid, which expresses gene products of PCV-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to the use of porcine circovirus-2 (PCV-2)
                                                                                                                                                                                                                                                                                                                        composition to treat pigs against myocarditis, abortion, intrauterine infection and/or post-weaning, multisystemic wasting syndrome associated with PCV-2
                                                                                                                                                                                                                                                                ATGACGTATCCAAGGAGGCGTTACCGAAGAAGAAGACACCGCCCCGCAGCCATCTTGGC
                                                                                                                                                                                                                                                                                       ATGACGTATCCAAGGAGGCGTTACCGCAGAAGAAGACACCGCCCCCGCAGCCATCTTGGC
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                                                                                                                                                                                                                   88.6%; Score 622; DB 22; Length 2520; 92.9%; Pred. No. 1e-177; O. Mismatches 50; Indels 0
                                  porcine circovirus-2 immunogen to formulate a vaccine
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                                                                                           2 #3; Fig 3 #1; 134pp; English
                                                                                                                                                                                                             Ouery Match 62.9
Best Local Similarity 92.9
WPI; 2001-244408/25.
P-PSDB; AAB73272.
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The present sequence was used in the construction of a recombinant virus comprising DNA from porcine circovirus 2 (PCV2). The recombinant virus is useful as vaccine for treatment and prophylaxis of PCV2 infection, such as postweaning multisystemic wasting syndrome (PMWS)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel recombinant virus comprising DNA from porcine circovirus 2 useful as vaccine for treatment and prophylaxis of porcine circovirus infection, such as postweaning multisystemic wasting syndrome in pigs
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                                                                                                                                                                                  PCV2; porcine circovirus 2; virucide; immunostimulant; vacci
postweaning multisystemic wasting syndrome; PMWS; infection;
pig pathogen; open reading frame 2; ORF2; ss.
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Pred. No. 1e-177;
^. Mismatches 50; Indels
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                                                                                                                                       pJP102 donor plasmid for PCV2 ORF2
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92.9%;
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01-JUN-2000; 2000US-0583545.
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                                                                                            (first entry)
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Matches 652; Conservative
  DNA;
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  standard;
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0; Mismatches
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CURRENT FILING DATE: 1999-03-12
EARLIER APPLICATION NUMBER: 60/077,890
EARLIER FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 3.0
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67.0%;
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SEQ ID NO 6
LENGTH: 1768
TYPE: DNA
ORGANISM: Porcine circovirus
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                                                                                                                                                                        Best Local Similarity 92.2
Matches 357; Conservative
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Matches 470; Conservative
                                                                             NAME/KEY: variation
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LENGTH: 69
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APPLICANT: HASSARD, LOZI
APPLICANT: HASSARD, John
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARPEYRE, Gilles E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
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                                                                                                                                                                                     Length 1768;
                                                                                                                                                                                                                30; Indels
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CURRENT APPLICATION NUMBER: US/09/161,092

CURRENT FILING DATE: 1998-09-25

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-20

PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-03

NUMBER OF SEQ. ID NOS: 6

SOFTWARE: PALENTIN VAF. 2.0
                                                                                                      NAME/KEY: variation
CCATION: (1)..(1768)
COTHER INFORMATION: N represents A or C or G or T
C-09-082-558-6
                                                                                                                                                                                  48.3%; Score 339; DB 4;
92.2%; Pred. No. 9.4e-96;
Live 0; Mismatches 30
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Patent No. 6391314
                                                         TYPE: DNA ORGANISM: Porcine circovirus
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CLARK, Edward
HAINES, Deborah
HASSARD, Lori
HARDING, John
              2.0
                                                                                                                                                                                                                  Matches 357; Conservative
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APPLICANT: ALLAN, Gordon
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                   Similarity
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                                            LENGTH: 1768
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                              SEQ ID NO 6
                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                   Best Local
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                                                                                             FEATURE:
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1528 AGGCTACAAACCTCTAGAAATGTGGACCACGTAGGCCTCGGCACTGCGTTCGAAAACACT 1469
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                                                                                                                                                                                                                               376 GATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTCC
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                                                                           Length 1768;
                                                                                                                Indels
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APPLICANT: Ritchie, Branson W.
APPLICANT: Niagro, Frank D.
APPLICANT: Lukert, Phil D.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines against Circovirus Infections FILE REFERENCE: 21099.0057
                                                                                                                30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 4;
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                                                                         Score 339; DB 4;
Pred. No. 9.4e-96;
0; Mismatches 30
                     or
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LOCATION: (1)..(1768)
OTHER INFORMATION: N represents A or C or G
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ογ	61	CAGATCCTCGGCGCGCGCCCTGGCTCGTCCACCCCGCCACGTTACCGCTGG 114
qa	61	61 AACATCCTCCGGAGAAGACCATATTTGGCACACCCCCTTCAGAACCGTTACAGATGG 120
Qy	115	115 AGAAGGAAAAATGGCATCTTCAACACCCGCCTCCCGCACCTTCGGATATACTGTCAAG 174
qa	121	CGCCGAAAGACGGGTATCTTCAATTCCCGCCTTTCTACAGAATTTGTACTCACCATAAAA 180
Qy	175	CGAACCACAGTCAGAACGCCCTCCTGGGCGGTGGACATGATGATTAATGAC 234
qa	181	181 GGAGGATACTCGCAGCCATCTTGGAATGTTAACTACCTCAAATTCAACATCGGCCAG 237
Qy .	235	TTTCTTCCCCCAGGAGGGGGGTCAAACCCCCGCTCTGTGCTGCCTTTGAATACTACAGAATA 294
Db	238	TTCCTCCCCCCTCAGGCGGCACCACCCCTACCCTTTCCAATACTACCGTATT 297
Qy	. 295	AGAAAGGTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTG 354
Db	298	AGAAAGGCTAAATGAATTTTACCCCAGAGACCCCATCACTCTAATCAAGAGGTGTT 357
Qy	355	GGCTCCAGTGCTGTTATTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTAT 414
qa	358	GGGTCCACTGTTGTTTTTTTTTGTACTTTGTAACCCCCTCCACTTGGCCTAT 417
Qy	415	GACCCCTATGTAAACTACTCCTCCCGCCATACCATAACCCAGCCCTTCTCTCTACCACCC 474
qq	418	GACCCCATATAATAACTACTCCTCCCGCCACATAAGGCAGCCCTTTACCTACC
Qy	475	475 CGGTACTTTACCCCCAAACCTGTCCTAGATTTCACTATTGATTACTTCCAACCAA
QQ	478	AGGTACTTCACCCCCAAACCTGACCTAACAATGATTGGTTCCACCAAATAAT 537
Oy .	535	
qa .	538	AAAAGAAACCAGCTGTGGCTCCATTTAAATACCCACACCATGTCGAGCACACAGGCCTC 597
Qy	595	GGCACTGCGTTCGAAAACAGTATACGACCAGGATACAATATCCGTGTAACCATGTAT 654
Db	598	GGCTATGCGCTCCAAAATGCAGCCACAGAGTTATGTGGTAAGGCTGACTATTAT 657
Qy	655	GTACAATTCAGAGAATTTAATTTTAAAGACCCCCACTTAA 695
qa	658	

Search completed: May 18, 2003, 12:47:57 Job time : 65 secs

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Best Local Sim
Matches 358;
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APPLICANT: MEEHAN, Brian
APPLICANT: MEEHAN, Edward
APPLICANT: CLARK, Edward
APPLICANT: HAINES, Deborah
APPLICANT: HASSARD, Lori
APPLICANT: HASSARD, Lori
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARREYRE, CATHERING TITLE OF INVENTION: REAGENTS
FILLE OF INVENTION: REAGENTS
FILLE REFERENCE: ALLAN
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Pred. No. 4.6e-96;
0; Mismatches 30
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CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: ER 9800873
EARLIER FILING DATE: 1998-01-22
EARLIER FILING DATE: 1998-01-22
EARLIER APPLICATION NUMBER: ER 9803707
EARLIER FILING DATE: 1998-03-20
EARLIER FILING DATE: 1998-03-03
EARLIER FILING DATE: 1998-03-03
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PALENTIN Ver. 2.0
TTTTAAAGACCCCCCACTTAACCCTTAA 702
                                                                                                                                                                                    Sequence 3, Application US/09082558A Patent No. 6368601 GENERAL INFORMATION:
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; ORGANISM: Porcine circovirus
US-09-082-558-3
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Matches 358; Conserva
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US-09-082-558-4/c
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1528 GAGACTACAAACCTCTGGAAATGTGGACCACGTAGGCCTCGGCGCTGCGTTCGAAAACAG 1469
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                                                               APPLICANT: HASSARD, Lori
APPLICANT: HASDING, John
APPLICANT: CHARREYE, Catherine E.
APPLICANT: CHAPPUIS, GILLes E.
APPLICANT: CHAPPUIS, GILLes E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
FILE REFERENCE: ALLAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 1768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 340; DB 4;
Pred. No. 4.6e-96;
0; Mismatches 30;
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                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/082,558A CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: FR 9800873
EARLIER FILING DATE: 1998-01-22
EARLIER FILING DATE: 1998-03-20
EARLIER FILING DATE: 1998-03-20
EARLIER FILING DATE: 1998-03-20
EARLIER FILING DATE: 1997-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48.4%; Score 340; 92.3%; Pred. No. 4
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Patent No. 6391314 GENERAL INFORMATION:
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HAINES, Deborah
HASSARD, Lori
HARDING, John
                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                   APPLICANT: CLARK, Edward APPLICANT: HAINES, Deborah
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APPLICANT: MEEHAN, Brian
MEEHAN, Brian
                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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LENGTH: 1768
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                                                 FILE REFERENCE: ALLAN
CURRENT APPLICATION NUMBER: US/09/161,092
CURRENT APPLICATION NUMBER: US/09/061,092
CURRENT FILING DATE: 1998-00-25
PRIOR APPLICATION NUMBER: EARLIER PAPLICATION NUMBER: 09/082,558
PRIOR FILING DATE: EARLIER FILING DATE: 1998-05-21
PRIOR FILING DATE: BARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: BARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: BARLIER FILING DATE: 1998-01-20
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-20
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 98/03707
PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 6
SEQ ID NOS: 6
SEQ ID NO 3
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APPLICANT: CLARK, Edward
APPLICANT: CLARK, Edward
APPLICANT: HAINES, Deborah
APPLICANT: HASSARD, Lori
APPLICANT: HARDING, John
APPLICANT: CHARRETRE, Catherine E.
APPLICANT: CHARRETRE, Catherine E.
TTILE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
APPLICANT: CHAPPUIS, Gilles E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
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Pred. No. 4.6e-96;
0; Mismatches 30
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CURRENT FILING DATE: 1998-09-25
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ilarity 92.3%;
Conservative
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APPLICANT: ALLAN, Gordon
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Best Local Simi
Matches 358; (
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US-09-161-092-4/c
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APPLICANT: ALLAN, GORDON
APPLICANT: ALLAN, GORDON
APPLICANT: CLARK, Edward
APPLICANT: CLARK, Edward
APPLICANT: HAINES, Deborah
APPLICANT: HASSARD, Lori
APPLICANT: HARDING, John
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARREYRE, Catherine E.
APPLICANT: CHARPINS, Gilles E.
APPLICANT: CHARPINS, GILLES E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
                                                                                                                        DATE: 1998-03-20
APPLICATION NUMBER: FR 97/12382
DATE: 1997-10-03
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                                                                                                   APPLICATION NUMBER: FR 9803707
       09/082,558
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       APPLICATION NUMBER:
                            DATE: 1998-05-21
APPLICATION NUMBER:
                                                                              DATE: 1998-01-22
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Pred. No. 4.6e-96;
0; Mismatches 30
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PRIOR APPLICATION NUMBER: EARLIER APPRIOR PRIOR DATE: EARLIER FILING DA PRIOR PRIOR FILING DATE: EARLIER FILING DA PRIOR FILING DATE: EARLIER FILING DA PRIOR APPLICATION NUMBER: EARLIER APPRIOR APPLICATION NUMBER: EARLIER APPRIOR APPLICATION NUMBER: EARLIER APPRIOR PILING DATE: EARLIER FILING DA NUMBER: EARLIER PILING DA NUMBER OF SEQ ID NOS: 6
SOFTWARE: PATCHIN VET. 2.0
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Best Local S:
Matches 358
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US-09-161-092-2/c
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APPLICANT:
APPLICANT:
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APPLICANT: ALLAN, Gordon
APPLICANT: CLARK, Edward
APPLICANT: CLARK, Edward
APPLICANT: CLARK, Edward
APPLICANT: HANSARD, Lori
APPLICANT: HASSARD, Lori
APPLICANT: CHAPPUIS, Catherine E.
APPLICANT: CHAPPUIS, Gilles E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
FILE REFERENCE: ALLAN
CURRENT APPLICATION NUMBER: US/09/082,558A
CURRENT FILING DATE: 1998-05-21
EARLIER RILING DATE: 1998-01-22
EARLIER FILING DATE: 1998-03-20
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        APPLICANT: KRAKOWKA, George S.
APPLICANT: KRAKOWKA, George S.
APPLICANT: AUDONNET, Jean-Christophe F.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08777
EARLIER FILING DATE: 1998-07-06
NUMBER: OF SEQ ID NOS: 5
SOFTWARE: PALENTIN VET. 2.1
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Pred. No. 4.9e-108;
0; Mismatches 6;
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                                                                                                                                                                                                                                                                                       Query Match 53.9%;
Best Local Similarity 98.5%;
Matches 382; Conservative
                                                                                                                                                                                                                                      ; ORGANISM: Porcine circovirus US-09-347-594-2
APPLICANT: ELLIS, John A.
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CURRENT FILING DATE: 1988-09-25
FRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/
PRIOR PELING DATE: EARLIER FILING DATE: 1998-05-21
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FLILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FLILING DATE: EARLIER FILING DATE: 1998-03-20
PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-20
PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-03
NUMBER: OF SEQ ID NOS: 6
SOFTWARE: PALENTING DATE: EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                               Score 378.4; DB 4;
Pred. No. 4.9e-108;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09161092 Patent No. 6391314
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
                                                                                                                                                                                               53.9%;
98.5%;
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APPLICANT: ALLAN, Gordon
APPLICANT: MEEHAN, Brian
APPLICANT: CLARK, Edward
APPLICANT: HAINES, Deborah
APPLICANT: HASSARD, LOOI
                                                                                                        TYPE: DNA ORGANISM: Porcine circovirus
                                                                                                                                                                                                                                          Matches 382; Conservative
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Best Local Similarity
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TGTTCTTGACTCCACTATTGATTACTTCCAACCAATAACAAAAAGGAATCAGCTTTGGCT 1529
                                                                                        555 GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG 614
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APPLICANT: ALLAN, GOCDON M.
APPLICANT: MELHAN, Brian M.
APPLICANT: MELLS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: AUDONNET, Jean-Christophe F.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08777
EARLIER FILING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
NUMBER OF SEQ ID NOS: 5
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Pred. No. 4.6e-96;
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Patent No. 6217883
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11 Similarity 92.3%;
358; Conservative
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SEO ID NO 4
LENGTH: 1768
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US-09-347-594-4/C
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APPLICANT: ALLAN, Gordon M.
APPLICANT: ALLAN, Brian M.
APPLICANT: MEEHAN, Brian M.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: KRAKOWKA, George S.
APPLICANT: KRAKOWKA, George S.
TILE REFERENCE: 454313-2338
CURRENT APPLICANTON: PORKINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08777
EARLIER APPLICATION NUMBER: 98 08777
EARLIER APPLICATION NUMBER: 98 08777
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN Ver. 2.1
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Pred. No. 4.9e-108;
0; Mismatches 6;
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Pred. No. 4.6e-96;
0; Mismatches 30,
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Patent No. 6217883
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Conservative
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Best Local Similarity 98.5%;
Matches 382; Conservative
; ORGANISM: Porcine circovirus US-09-161-092-2
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4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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APPLICANT: MERHAN, Brian
APPLICANT: MERHAN, Brian
APPLICANT: CLARK, Edward
APPLICANT: HANDING, Deborah
APPLICANT: HANDING, John
APPLICANT: GRAREFYER, Catherine E.
APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: CHAPPUIS, GILLS F.
TITLE OF INVENTION: REAGENTS
FILE REFERENCE: ALLAN
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Pred. No. 5e-109;
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CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: FR 9800873
EARLIER FILING DATE: 1998-01-22
EARLIER APPLICATION NUMBER: FR 9803707
EARLIER FILING DATE: 1998-03-20
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 6
SEQ ID NO 1
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99.0%;
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RESULT 3

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                                                                                                                                                                                                         APPLICANT: CHARRETR, Catherine E.
APPLICANT: CHARRETR, Catherine E.
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
FILE REFERENCE: ALLAN
CURRENT PAPLICATION NUMBER: US/09/161,092
CURRENT FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: O9/082,558
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-32
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-20
PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-003
NUMBER OF SEQ ID NOS: 6
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Pred. No. 5e-109;
0; Mismatches 4;
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Sequence 1, Application US/09161092
Patent No. 6391314
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99.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Porcine circovirus
                                                                                          MEEHAN, Brian
CLARK, Edward
HAINES, Deborah
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Matches 384; Conservative
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                                                                                                                                                              HASSARD, Lori
HARDING, John
                                                                         Gordon
                                              GENERAL INFORMATION:
                                                                         APPLICANT: ALLAN,
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fruit fly.
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Matches 56;
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Determination of this BAC-end sequence was carried out as part of Determination of this BAC-end sequence was carried out as part of Collaboration with the Berkeley Drosophila Genome Project (BDGP).
The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC ilbrary was prepared by Kazutoyo Gsoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial EcoRI digestion of Drosophila DNA provided by the BDGP from the isogenic strain v2; on bw sp. the same strain used for the BDGP's Pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be Location/Qualifiers

Location/Qualifiers

Location/Qualifiers
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                                                                        /Gev_mcr________/dev_steage="adult"
/lab_host="DHS - alpha"
/lab_host="DHS - alpha"
/note="Organ: head: Vector: pOT2; Site_1: EcoRI; Site_2:
XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
                                                                                                                                                                                                                                                                                                                                                        Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                       32 GAAGACACCGCCCCCGCAGCCATCTTGGCCAGATCCTCCGCCGCCGCCCCTGGCTCGTCC
                                                                                                                                                                                                                                                                                                                                                                                                               92 ACCCCCCCCACCGTTACCGCTGGAGAAGGAAAATGGCATCTTCAACACCCGCCTCTCCC
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Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilae.
1 (bases 1 to 939)
                                          /clone_lib="GH Drosophila melanogaster head pOT2
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Pred. No. 0.12;
0; Mismatches 56; Indels
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/db_xref="taxon:7227"
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                                                             /sex="male and female"
/db_xref="taxon:7227"
/clone="GH27112"
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/note="end : TET3"
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Bukaryotta; Metazoa: Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 638)
Stapleton,M., Brokstein,P., Hong,L., Tyler,D., Berman,B., Carlson,J., Champe,M., Chavez,C., Dorsett,V., Farfan,D., Frise,E., George,M., Gonzalez,M., Guarin,H., Harris,N., Li,P., Liao,G., Misra,S., Mungall,C.J., Nunoo,J., Pacleb,J., Paragas,V., Park,S., Phouanenavong,S., Wan,K., Yu,C., Lewis,S.E., Celniker,S. and Rubin
                                                                                                                                                   391 ACAAAGGCCACACCCTCACCTATGACCCCTATGTAAACTACTCCTCCCGCCATACCATA 450
                                                                                                                                                                                              274 MMMHAWMCCTTCCCTCCCYCCYTCCCCCCAYYYMCCCMCCCTTCTYCCCATCCMMC 333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="RH Drosophila melanogaster normalized Head
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hit genomic AE003629; arm:2L [10413221,10674011]
estimated-cyto:31E5-3255:08/23/2001
Plate: RH.583 row. A column: 10
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          Length 939;
6.4%: Score 44.6; DB 17; Length ilarity 18.1%; Pred. No. 0.23; Conservative 122; Mismatches 131; Indels
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/db_xref="taxon:7227"
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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Scoring table:

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A1238309 GH14332.5
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A1238565 GH14632.5
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AL151849 Anopheles
AM015354 Hh74b04.x
AI141028 oy68h12.x
AI364938 q223402.x
BI799844 H142B04 E
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Eukaryota: Metazoa: Arthropoda: Hexapoda; Insecta; Pterygota;
Eukaryota: Endopterygota: Diptera; Brachycera; Muscomorpha:
Ephydroidea: Drosophilidae; Drosophila.
1 (bases 1 to 561)
Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang,G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 271 row: A column: 12
High quality sequence stop: 488.
Location/Qualifiers
1 .561
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
                                                                                                                                                                                                                                                                                                                                                                                                                                      ALIGNMENTS
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BDGP/HHMI Drosophila EST Project
Unpublished (2001)
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AI514058.1 GI:4418120
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BI629442 RH58310.5
AL292664 GH15617.5
AL108264 GH07058.5
AL405757 GH25879.5
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                                                                        May 18, 2003, 10:32:44; Search time 1499 Seconds (without alignments) 7584.544 Million cell updates/sec
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            GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                           16154066 segs, 8097743376 residues
                                                                                                                                                                                                              Potal number of hits satisfying chosen parameters:
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                                                       nucleic search, using sw model
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CNS00CNG
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AI108264
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Maximum Match 100%
Listing first 45 su
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gb_htc:*
gb_est3:*
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C99980 C99980 Oryz

BO704127 946110E10 BH879805 ht48h11.9

BG599848 EST504743 BM110666 EST558202

AA698799 HL05714.5

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fruit fly.
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/dev_stage="adult"
/lab_nost="DHs - alpha"
/note="Yorgan: head; Vector: pOT2; Site_1: EcoR1; Site_2:
Xho1; Sized fractionated cDNAs were directly ligated into
pOT2. Plasmid cDNA library."
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophiladae; Drosophila.
I (bases I to 47)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G., Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
BDGP
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                                                                                            Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Mosptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 493)
1 (bases 1 to 493)
1 Larvey, D., Brokstein, P., Hong, L., Evans-Holm, M.; Su, C., Tsang, G. BDGP/HHMI Drosophila EST Project
                                                                                                                                                                                                                                                                                    Lawrence Berkeley National Lab One Cyclotron Rd, Berkeley, CA 94720, USA Exa: 510 486 6798 Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu Plate: 238 row: G column: 7 High quality sequence stop: 492. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Drosophila melanogaster"
/db_xref="taxon:7227"
/clone="GH25879"
/clone_lib="GH Drosophila melanogaster head pOT2"
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Pred. No. 0.29;
0; Mismatches 55; Indels
                                                                          Drosophila melanogaster
                                                                                                                                                                                                                             Unpublished (2001)
Contact: Stapleton, M.
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                 AI405757.1 GI:4248844
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AII34557 532 bp mRNA linear EST 19-APR-2001 GH12051.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH12051 5prime, mRNA sequence.
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/lab.host="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 532)
1 (bases 1 to 532)
1 Lewis, S. and Rubin, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,
Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
                                                                                   Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 143 row: C column: 8
High quality sequence stop: 457.
Location/Qualiflers
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Plate: 120 row: E column: 3
High quality sequence stop: 403.
Location/Qualifiers
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/dev_stage="ballt" alpha"
                                                                                                                                                                                                                                                                                                                       /clone_lib="GH Drosophila melanogaster head pOT2"
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/dev_stage="adult"
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/db_xref="taxon:7227"
/clone="GH14332"
Lawrence Berkeley National Lab
One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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Fax: 510 486 6798
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Matches 77; Conservative
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Matches 77;
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/sex="male and female"
/dev_stage="adult"
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/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2:
XhoI; Sized fractionated cDNAs were directly ligated into
                                                                                                                                                                                                                                                                        TCCACCCCCCCCCCCTCCCTTGCCCTTGGGAAGGAAAAATGGCATCTTCAACACCCCCTCT 148
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                                                                                                                                                                                                                                       581 GCNGCCGATTCCACCGCCTCCACCAGAATGGCCACCTCCGCCGCCGATTCCGCCACAAA 522
/lab_host="DH5-alpha TonA"
/note="Organ: head; Vector: pFlc1; Site_1: XhoI; Site_2:
Monte="Organ: head; Vector: pFlc1; Site_1: XhoI; Site_2:
BamHI; Library was kindly generated by Piero Carninci at
the RIKEN. The library was normalized and excised using
Cre recombinase. Plasmid cDNA library."
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Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
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Plate: 156 row: B column: 5
High quality sequence stop: 293.
Location/Qualifiers
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Pred. No. 0.28;
0; Mismatches 55; Indels
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    .445
/organism="Drosophila melanogaster"
/db_xref="taxon:7227"

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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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1 (bases 1 to 445)
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                                                                                                                                     DB
                                                                                                                                    Score 44.2; DB Pred. No. 0.27; 0; Mismatches
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Similarity 57.2%;
79; Conservative
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AI108264 484 bp mRNA linear EST 19-APR-2001 GH07058.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH07058 5prime, mRNA sequence.
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/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: XhoI; Sized fractionated cDNAs were directly ligated into
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95 CCCGCCACCGTTACCGCTGGAGGAAAAATGGCATCTTCAACACCCGCCTCTCCCGCA 154
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( (bases 1 to 484)
Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang, Lewis,S. and Rubin,G.M.
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Plate: 70 row: E column: 10
High quality sequence stop: 404.
Location/Qualifiers
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/sex="male and female"
/dev_stage="adult"
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Pred. No. 0.29;
0; Mismatches 55; Indels
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/db_xref="taxon:7227"
/clone="GH07058"
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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Drosophila melanogaster
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Query Match 6.3%;
Best Local Similarity 58.3%;
Matches 77; Conservative
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    Fax: 510 486 6798
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/dev_stage="adult"
/lab_host="adult"
/lab_host="organ: head; Vector: pOT2; Site_1: EcoRI; Site_2:
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2:
XNoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
109 c 241 g 133 t
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                                  Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;

Eukaryota; Endopterygota; Diptera; Brachycera; Muscomorpha;

Ephydroidea; Drosophilidae; Drosophila.

I (bases 1 to 558)

Lavrey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang,G.,

Lewis,S. and Rublin,G.M.

BDGP/HHMI Drosophila EST Project

Unpublished (2001)
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 466 6798
Email: http://www.fruitfly.org/EST, estefruitfly.berkeley.edu
Plate: 146 row: C column: 8
High quality sequence stop: 489.
Location/Qualifiers
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 561)
                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Drosophila melanogaster"
/db xref="taxon:7227"
/clone="6114632"
/clone=11b="GH Drosophila melanogaster head pOT2"
/sex="male and female"
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Pred. No. 0.3;
0; Mismatches 55; Indels
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One Cyclotron Rd, Berkeley, CA 94720, USA
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Unpublished (2001)
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AI404901.1 GI:4247988
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Best Local Similarity 58.3%;
Matches 77; Conservative
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AUTHORS
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Ada_xref="taxon:7227"

Ada_xref="taxon:7227"

Aclone="GH24758"

Aclone="GH24758"

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Aclone="taxon:727"

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1 (bases 1 to 561)
Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang,G., Ebais,S. and Rubin,G.M.

BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 247 row: E column: 10
High quality sequence stop: 461.
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, estefruitfly.berkeley.edu
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/sex="male and female"
/dev_stage="adult"
/lab_host="DH5 - alpha"
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/db_xref="taxon:7227"
/clone="GH26478"
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Pred. No. 0.3;
0; Mismatches
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High quality sequence stop: 494.
Location/Qualifiers
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AI406248.1 GI:4249335
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AI406114 557 bp mRNA linear EST 19-APR-2001 GH26313.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH26313 5prime, mRNA sequence.
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1 (bases 1 to 557)

1 (bases 1 to 557)

Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, Lewis, S. and Rubin, G.M.

BDGP/HHMI Drosophila EST Project
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plate: 263 row: B column: 1
High quality sequence stop: 494.
Location/Qualifiers
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/db_xref="taxon:7227"
/clone="cH26313"
/clone_lib="GH Drosophila melanogaster head pOT2"
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58.3%; Pred. No. 0.3;
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Fax: 510 486 6798
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/lab_host="DH5 - alpha"
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AI238565/c
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  /note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
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[ (bases I to 557)

Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G., Lewis, S. and Rubin, G.M.

BDGP/HHMI Drosophila EST Project
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Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 237 row: A column: 5
High quality sequence stop: 455.
Location/Qualifiers
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/sex="male and female"
/dev_stage="adult"
/lab_host="DH5 - alpha"
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                                                                                                                         Length 532;
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/organism="Drosophila melanogaster"
/db_xref="taxon:7227"
/clone="GH23705"
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                                                                                                                           Score 44; DB 9;
Pred. No. 0.29;
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ilarity 58.3%;
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Drosophila melanogaster

Eukaryota; Metazoa; Arhropoda; Hexapoda; Insecta; Pterygota;

Neoptera; Medazoa; Diptera; Brachycera; Muscomorpha;

Ephydroides; Drosophilidae; Drosophila.

I (bases 1 to 575)

Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,

Lewis, S. and Rubin, G.M.

Body/Hant Drosophila EST Project

Unpublished (2001)

Contact: Stapleton, M.
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Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 226 row: A column: 2
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/organism="Drosophila melanogaster"
/db_xref="taxon:727"
/dlone="6H22602"
/clone=lib="GH Drosophila melanogaster head pOT2"
/sex="male and female"
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                                                                                                                                  DB 9; Length 561;
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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Pred. No.
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High quality sequence stop: 437.
Location/Qualifiers
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Matches 77; Conserv
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AI516823 593 bp mRNA 11near EST 19-APR-2001 GH27392.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH27392 5prime, mRNA sequence.
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BDGP/HHMI Drosophila EST Project
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 273 row: H column: 8
High quality sequence stop: 547.
Location/Qualifiers
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Eukaryota: Metazoa: Arthropoda: Hexapoda: Insecta: Pterygota:
Neoptera: Endopterygota: Diptera: Brachycera: Muscomorpha:
Ephydroidea: Drosophilidae: Drosophila
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Pred. No. 0.3;
0; Mismatches 55; Indels
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
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AI516823.1 GI:4419923
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                              | || | || || 437 TCCTCCGGAATA 426
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Matches 77; Conservative
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1735 ATGACGTATCCAAGGAGGCGTTACCGCAGAAGAAGACACCGCCCCCGCAGCCATCTTGGC 1676
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APPLICANT: WANG, LI
APPLICANT: WANG, LI
APPLICANT: BABIUK, LORNE A.
APPLICANT: POTTER, ANDREW A.
APPLICANT: WILLSON, PHILIP
TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: 2001-08-20
CURRENT APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/209,961
PRIOR PLILING DATE: EARLIER FILING DATE: 1997-12-11
PRIOR PLILING DATE: EARLIER FILING DATE: 1997-12-11
PRIOR PLILING DATE: EARLIER FILING DATE: 1997-12-16
NUMBER: PARLIER FILING DATE: EARLIER FILING DATE: 1997-12-16
NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-16
                                                     GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCC
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SEQ ID NO 24
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                                                 CCCCCAGGAGGGGGGTCAAACCCCCGCTCTGTGCCCTTTGAATACTACAGAATAAGAAAG
                                                                                                                                                                                                                                                 Sequence 1, Application US/09935428A
Betent No. US20020106639A1
GENERAL INFORMATION:
APPLICANT: WANG, LI
APPLICANT: DOTTER, ANDREW A.
APPLICANT: WILLSON, PHILIP
TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
                                                                                                                                                                  GTTAAGGTTGAATTCTGGCCCTGCTCCCCGGATCACCCAGGGTGACAGGGGAGTGGGGCTCC
                                                                                                                                                                                                                                                                                                   TATGTAAACTACTCCTCCGGCCATACCATAACCCAGCCCTTCTCCTACCACTCCGGGTAC
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CURRENT APPLICATION NUMBER: US/09/935,428A
CURRENT APPLICATION NUMBER: US/09/935,428A
CURRENT FILING DATE: 2001-08-20
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/PRIOR FILING DATE: 1998-12-10
PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-11
PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-11
PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-16
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PATCHTIN VET. 2.0
SEQ ID NO
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Pred. No. 2.3e-195;
0; Mismatches 51;
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JS-09-935-428A-1
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ilarity 92.7%;
Conservative (
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Matches 651; Conserv
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US-09-935-428A-1/c
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1256

480

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Sequence 2, Appli
Sequence 3, Appli
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Sequence 1, Appl
                                                                                            (without alignments)
5521.565 Million cell updates/sec
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1 atgacgtatccaaggaggcg.....accccccacttaacccttaa 702
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                                                                                                                                                                                                                                                                                                                                                                                             1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
8: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
10: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
11: /cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
12: /cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
12: /cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
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                                                                                ; Search time 164 Seconds
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GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-935-428A-1
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US-10-112-540-1
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US-09-884-514-1
US-09-884-514-2
US-09-784-962-2
US-09-784-962-4
US-09-784-514-3
US-09-884-514-3
US-09-884-514-6
US-09-935-4288-5
US-09-935-4288-5
US-09-038-001-7
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US-09-884-514-5
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Maximum Match 100%
Listing first 45 summaries
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                                                      nucleic search, using sw model
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length: 2000000000
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sequence 466, App Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 54, Appl Sequence 332, App Sequence 332, App Sequence 1058, Ap Sequence 48, Appl Sequence 3, Appl Sequence 3, Appl Sequence 1921, Ap Sequence 5, Appli Sequence 3, Appli Sequence 1, Appli Sequence 2102, Ap Sequence 2300, Ap Sequence 239, App Appl Sequence 24, 0 US-09-802-669-24 0 US-09-801-107-2102 0 US-09-80-107-2300 0 US-09-92-254-239 0 US-09-92-24-88 0 US-10-115-695-48 0 US-10-115-61-48 0 US-10-115-61-48 0 US-10-115-61-48 0 US-10-115-61-48 0 US-10-115-61-48 0 US-10-115-61-48 0 US-10-115-61-3 0 US-09-920-671-31 0 US-09-920-671-31 0 US-09-920-671-31 US-10-108-605-54 US-10-184-644-332 US-10-184-634-332 US-10-025-380-1058 US-09-784-962-1 US-09-884-5 2038 3715 13715 2462 2462 2462 2462 2462 13925 13925 1107 127197 1767 336.44 366.44 366.44 366.22 368.38 368.38

ALIGNMENTS

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1675 CAGATCCTCCGCCGCCCCTGGCTCGTCCACCCCGCCACCGCTACCGTTGGAGAAGG 1616
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                                                                                                GENERAL INFORMATION:
APPLICANT: WANG, LI
APPLICANT: WANG, LI
APPLICANT: BABIUK, LORNE A.
APPLICANT: POTTER, ANDREW A.
APPLICANT: WILLSON, PHILIP
TITLE OF INVENTION: POSTWBANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWBANING WILLISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWBANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWBANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: POSTWBANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION NUMBER: GARLIER APPLICATION NUMBER: 09/209,961
PRIOR PELLING DATE: EARLIER FILLING DATE: 1997-12-11
PRIOR APPLICATION NUMBER: EARLIER FILLING DATE: 1997-12-11
PRIOR APPLICATION NUMBER: EARLIER FILLING DATE: 1997-12-11
PRIOR APPLICATION NUMBER: EARLIER FILLING DATE: 1997-12-11
SPRIOR FILLING DATE: EARLIER FILLING DATE: 1997-12-11
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Pred. No. 2e-196;
0; Mismatches 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Porcine Circovirus Type II
                                                    Sequence 11, Application US/09935428A Patent No. US20020106639A1
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93.0%;
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                          US-09-935-428A-11/c
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Best Local Simi
Matches 653; (
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RESULT 1
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RESULT 7
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                             US-09-884-514-1
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APPLICANT: HASARD LOTI
APPLICANT: HARDING, John
APPLICANT: CHARREYE, Catherine E.
APPLICANT: CHARPEYE, Catherine E.
APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: NEWTONREDS, Francis McNeilly
TITLE OF INVENTION: REACENTS
TITLE OF INVENTION: REACENTS
TITLE OF INVENTION: REACENTS
TITLE OF INVENTION NUMBER: US/09/884,514
CURRENT FILING DATE: 2000-06-19
PRIOR APPLICATION NUMBER: FR 9800873
PRIOR FILING DATE: 1998-01-22
PRIOR FILING DATE: 1998-01-22
PRIOR FILING DATE: 1998-01-03
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PATENTIN VOIL 2.0
SOFTWARE: PATENTIN VOIL 2.0
SOFTWARE: PATENTIN VOIL 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                               315 CTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTTT
                                                                                                                                                                                                                                                 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC
                                                                                                                                                                                                                                                                                                      CTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTACTTTACCCCCAAACC
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                                                                                                                                                                           Gaps
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                                                                                                                                              Length 1767
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                                                                                                                                             Score 381.6; DB 10;
Pred. No. 5.2e-116;
0; Mismatches 4;
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              08777
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PRIOR FILING DATE: 1999-07-04
PRIOR APPLICATION NUMBER: 98 08:
PRIOR FILING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN VEr: 2.1
SEQ ID NO 1
                                                                                                                                              54.4%;
99.0%;
                                                                                        TYPE: DNA ORGANISM: Porcine circovirus
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CLARK, Edward
HAINES, Deborah
                                                                                                                                                                     Matches 384; Conservative
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APPLICANT: ALLAN, Gordon
                                                                                                                                                          Similarity
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                                                                             LENGTH: 1767
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                                                                                                                                          375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCTATGTAAACTACTACTC
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                                                            Length 1767;
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TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REPERENCE: 454313-2338
CURRENT ABLICATION NUMBER: US/09/784,962
CURRENT FILING DATE: 2001-02-16
PRIOR APPLICATION NUMBER: 09/347,594
PRIOR FILING DATE: 1999-07-04
PRIOR FILING DATE: 1999-07-06
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Pred. No. 6e-115;
0; Mismatches 6; Indels
                                                                                                    4; Indels
                                                          Score 381.6; DB 10
Pred. No. 5.2e-116;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           675 TTTTAAAGACCCCCCACTTAACCCTTAA 702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEEHAN, Brian M.
ELLIS, John A.
KRAKOWKA, George S.
AUDONNET, Jean-Christophe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09784962 Patent No. US20020146431A1
                                                            54.4%;
99.0%;
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Best Local Similarity 98.5%;
Matches 382; Conservative
ORGANISM: Porcine circovirus
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                                                        Query Match
Best Local Similarity 99.0
Matches 384; Conservative
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APPLICANT: ALLAN, Gordon M.
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; LENGTH: 1768
; TYPE: DNA
; ORGANISM: PO:
US-10-112-540-1
 LENGTH: 1768
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                 Length 1343;
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APPLICANT: Liu, Oiang
APPLICANT: Liu, Oiang
APPLICANT: TIKOO, Suresh K.
APPLICANT: Wilson, Philip
APPLICANT: Babiuk, Lorne A.
TITLE OF INVENTION: METHODS TO CULTURE CIRCOVIRUS
FILE REPERENCE: 293102003100;
CURRENT APPLICATION NUMBER: US/10/112,540
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: US 60/279,173
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ 1D NOS: 3
SOFTWARE: FastSEQ for Windows Version 4.0
                Score 618.8; DB 10;
Pred. No. 6.6e-195;
                                  52;
                                   0; Mismatches
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                88.1%;
92.6%;
                          Best Local Similarity 92.6
Matches 650; Conservative
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US-10-112-540-1/c
JS-09-935-428A-24
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                                                   Score 618.8; DB 9;
Pred. No. 7.7e-195;
                                                                                      52;
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APPLICANT: ALLAN, Gordon M.
APPLICANT: ELLIS, John A.
APPLICANT: REAKOWKA, George S.
APPLICANT: KRKOWKA, George S.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARF.
FILE REFERENCE: 454313-2338
CURRENT APPLICATION NUMBER: US/09/784,962
CURRENT FILING DATE: 2001-02-16
PRIOR APPLICATION NUMBER: 09/347,594
                                                                                    0; Mismatches
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Patent No. US20020146431A1
                                                   88.1%;
92.6%;
circovirus
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Porcine
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US-09-784-962-1/C
                                                   Query Match
Best Local S:
Matches 650
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Similarity
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LENGTH: 1768
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                                                                           APPLICANT: MEBHAN, BITAIN M.
APPLICANT: MEBHAN, BITAIN M.
APPLICANT: KRAKOWRA, GEORGE S.
APPLICANT: RANGOWRA, GEORGE S.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT FILING DATE: 2001-02-16
PRIOR FILING DATE: 2001-02-16
PRIOR FILING DATE: 1999-07-04
PRIOR APPLICATION NUMBER: 98 08777
PRIOR APPLICATION NUMBER: 98 08777
PRIOR APPLICATION NUMBER: 98 08777
PRIOR PELING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
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Pred. No. 3.5e-102;
0; Mismatches 30;
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                          Sequence 4, Application US/09784962
Patent No. US20020146431A1
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Patent No. US20020146432A1
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.larity 92.3%;
Conservative C
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; ORGANISM: Porcine circovirus
US-09-784-962-4
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APPLICANT: MERHAN, Brian
APPLICANT: CLARK, Edward
APPLICANT: HAINES, Deborah
APPLICANT: HASSARD, Lori
APPLICANT: HARBING, John
APPLICANT: CHARREYRE, Cather
                                                    GENERAL INFORMATION:
APPLICANT: ALLAN, Gordon M.
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358; Conserv
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RESULT 10
US-09-784-962-4/c
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Best Local S:
Matches 358
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APPLICANT: CHAPPUIS, Gilles E.
APPLICANT: NEWTONARDS, Francis McNeilly
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
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APPLICANT: HASSARD, Lori
APPLICANT: HASSARD, Lori
APPLICANT: CHARREYER, Catherine E.
APPLICANT: CHARREYER, Catherine E.
APPLICANT: NEWTONARDS, Francis McNeilly
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; DB 10;
3.5e-102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 340; DB
Pred. No. 3.5e-
0; Mismatches
                                                                                                     FILE REFERENCE: ALLAN
CURRENT APPLICATION NUMBER: US/09/884,514
CURRENT APPLICATION NUMBER: EN 9800873
PRIOR PILING DATE: 1998-01-22
PRIOR APPLICATION NUMBER: FR 9803707
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: FR 97/12382
PRIOR FILING DATE: 1999-03-20
PRIOR FILING DATE: 1997-10-03
NUMBER OF SEO ID NOS: 6
SOFTWARE: Patentin Ver: 2.0
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92.38;
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Matches 358; Conservative
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APPLICANT: MEEHAN, Brian
APPLICANT: CLARK, Edward
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1408 TCTTAAAGACCCCCACTTAAACCCTAA 1381
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US-09-784-962-3/c
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APPLICANT: ALLAN, WOLUGUE
APPLICANT: ALLAN, WOLUGUE
APPLICANT: HAINES, Deborah
APPLICANT: HAINES, Deborah
APPLICANT: HASARD, Lori
APPLICANT: HARDING, John
APPLICANT: CHARREYE, Catherine E.
APPLICANT: CHARREYE, Catherine E.
APPLICANT: CHARREYE, Catherine E.
APPLICANT: CHARREYE, Catherine E.
APPLICANT: RAPPLICANT: REAGENTS
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
FILE REFERENCE: ALLAN
CURRENT APPLICATION NUMBER: FR 9800873
PRIOR FILING DATE: 1998-01-22
PRIOR APPLICATION NUMBER: FR 9903707
PRIOR APPLICATION NUMBER: FR 97/12382
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: FR 97/12382
NUMBER OF SEQ ID NOS: 6
SSOTID NO 2
TENANTH: 1767
                           CTCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTACTTTACCCCCAAACC
                                                                               GAGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAG
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Pred. No. 6e-115;
0; Mismatches 6; Indels
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Best Local Similarity 98.5%;
Matches 382; Conservative
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APPLICANT: ALLAN, Gordon
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APPLICANT: ALLAN, Gordon M.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: AUDONNET, Jean-Christophe F.
TITLE OF INVENTION: POOFCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT FILLING DATE: 10909-07-04
PRIOR FILLING DATE: 1999-07-04
PRIOR FILLING DATE: 1999-07-04
PRIOR FILLING DATE: 1999-07-06
NUMBER OF SED ID NOS: 5
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Pred. No. 3.5e-102;
0; Mismatches 30;
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Best Local Similarity 92.33
Matches 358; Conservative
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 Pred. No. 1.7e-91;
0; Mismatches 222; Indels
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TITLE OF INVENTION: ROLLING CIRCLE REPLICON
TITLE OF INVENTION: VECTORS
TITLE REFERENCE: 008010179CPUS01
CURRENT APPLICATION NUMBER: US/10/038,001
CURRENT APPLICATION NUMBER: 09/505,477
PRIOR FILING DATE: 2000-02-16
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SOFTWARE: FastSEQ for Windows Version 4.0
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Publication No. US20020187952A1
GENERAL INFORMATION:
APPLICANT: POGGUE, Kenneth E.
APPLICANT: POGGUE, Gregory P.
APPLICANT: MCCORMICK, Alison
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 67.08;
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Best Local Similarity 67.0 Matches 470; Conservative
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US-10-038-001-7/c
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LENGTH: 7460
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 Length 7460;
 DB 9;
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Score 273.4; DB 9
Pred. No. 1.1e-79;
0; Mismatches 181
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Job time : 170 secs
 38.9%;
68.4%;
 Query Match 38.9
Best Local Similarity 68.4
Matches 411; Conservative
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PatentIn Ver.
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 SOFTWARE: Pater
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LENGTH: 1768
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US-09-935-428A-2
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TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
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CURRENT APPLICATION NUMBER: US/09/884,514
CURRENT FILING DATE: 2000-06-19
PRIOR APPLICATION NUMBER: FR 9803707
PRIOR PELICATION NUMBER: FR 9803707
PRIOR PELICATION NUMBER: FR 9803707
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: FR 97/12382
PRIOR FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    675 TTTTAAAGACCCCCCCACTTAACCCTTAA 702
PRIOR APPLICATION NUMBER: FR 9800873
PRIOR FILING DATE: 1998-01-22
PRIOR APPLICATION NUMBER: FR 9803707
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 4
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                                                                                                                                                               ORGANISM: Porcine circovirus
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CLARK, Edward
HAINES, Deborah
HASSARD, Lori
HARDING, John
                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                        Similarity
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1168 TGGCCCTGCTCCCCCATCACCCAGGTGATAGGGGAGTGGGCTCCACTGTTATTCTA 1709
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TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM
TITLE OF INVENTION: PIGS

TITLE OF INVENTION: PIGS

FILE REFERENCE: 9000-0040

CURRENT APPLICATION NUMBER: US/09/935,428A

CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/209,961

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069,233

PRIOR FILING DATE: BARLIER FILING DATE: 1997-12-11

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069,750

PRIOR FILING DATE: BARLIER FILING DATE: 1997-12-16

NUMBER OF SEQ ID NOS: 24

SEQ ID NO 2.20
                                                                                                                                                                                                                                                                        316 TGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTGGGCTCCAGTGCTGTTATTA 375
                                                                                                                                                                                                                                                                                                                                                                        376 GATGATAACTTTGTAACAAAGGCCACACACCCTCACCTATGACCCCTATGTAAACTACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         436 TCCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCGGTACTTTACCCCCAAACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       556 AGACTACAAACTGCTGGAAATGTAGACCACGTAGGCCTCGGCACTGCGTTCGAAAACAGT
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                                                                                                                                                                       Length 1768;
                                                                                                                                                                                                                          Indels
                                                                                                                                                                       Score 339; DB 10;
Pred. No. 7.5e-102;
0; Mismatches 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      676 TTTAAAGACCCCCCACTTAACCCTTAA 702
                                                                  LOCATION: (1)...(1768)
COTHER INFORMATION: N represents A or C US-09-884-514-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09935428A Patent No. US20020106639A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Porcine Circovirus Type
                                                                                                                                                                          48.3%;
92.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: WANG, LI
APPLICANT: BABIUK, LORNE A.
APPLICANT: POTTER, ANDREW A.
ORGANISM: Porcine circovirus
                                                                                                                                                                          Query Match
Best Local Similarity 92.2
Matches 357; Conservative
                                                  NAME/KEY: variation
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DB 10;

Score 307.8;

43.88;